

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 09/486,247B
Source: IFW/6
Date Processed by STIC: 5/20/05

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IFW16

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/486,247B

DATE: 05/20/2005
TIME: 09:07:27

Input Set : A:\31304-703.831.ST25.txt
Output Set: N:\CRF4\05202005\I486247B.raw

3 <110> APPLICANT: Deutsches Krebsforschungszentrum Stiftung des öffentlichen
 4 Rechts
 5 DEAR, Terence N.
 6 BOEHM, Thomas
 8 <120> TITLE OF INVENTION: PROTEASE-RELATED PROTEIN
 10 <130> FILE REFERENCE: 31304-703.831
 12 <140> CURRENT APPLICATION NUMBER: 09/486,247B
 13 <141> CURRENT FILING DATE: 2000-05-25
 15 <160> NUMBER OF SEQ ID NOS: 8
 17 <170> SOFTWARE: PatentIn version 3.2
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 822
 21 <212> TYPE: DNA
 22 <213> ORGANISM: Mouse
 24 <400> SEQUENCE: 1
 25 taggtggtgt cattcccttc caacctgagt gctggcaggt acactgctgg ccaccagcag 60
 27 atgcccattga agatgctgac aatgaagatg ctggccctgt gtttgcgttct tgctaaatca 120
 29 gcctggtcgg aggaacagga gaaggtgggt catggaggcc cgtgtttgaa ggactcccac 180
 31 ccttccagg ctgcctcta cacctcagggt cacttgcgtgt gtgggtgggt cctcattgac 240
 33 ccacagtggg tgctgacagc tgcccactgc aaaaaaccga atctgcaggat gatcttgggg 300
 35 aaacacaacc tacggcaaacc agagactttc caaaggcaaa tctcaggatggc caggactatt 360
 37 gtccatcccc gctacaaccc taaaacccac gacaatgaca tcataatgtgt gcatctgaaa 420
 39 aatccagtca aattctctaa aaagatccag cctctgcctt tgaagaatga ctgcctgag 480
 41 gagaatccca actgccagat cctgggctgg ggcaagatgg aaaatggtga cttcccgat 540
 43 accattcagt gtgtgtatgt ccatctgggt cccggggagc agtgcgtggc tgcctaccct 600
 45 ggcagatca cccagagcat ggtgtgcgc ggcacatga aagaaggcaa cgatccctgt 660
 47 cagggtgatt ctggaggtcc cctagtatgt ggggtcgcc tccgagggtct cgtgtcatgg 720
 49 ggtgacatgc cctgtggatc aaaggagaag ccaggagttt acaccgtatgt ctgcactcat 780
 51 atcagatggc tccaaacat cctcagaaac aagtggctgt ga 822
 54 <210> SEQ ID NO: 2
 55 <211> LENGTH: 253
 56 <212> TYPE: PRT
 57 <213> ORGANISM: Mouse
 59 <400> SEQUENCE: 2
 61 Met Pro Met Lys Met Leu Thr Met Lys Met Leu Ala Leu Cys Leu Val
 62 1 5 10 15
 65 Leu Ala Lys Ser Ala Trp Ser Glu Glu Gln Glu Lys Val Val His Gly
 66 20 25 30
 69 Gly Pro Cys Leu Lys Asp Ser His Pro Phe Gln Ala Ala Tyr Thr
 70 35 40 45
 73 Ser Gly His Leu Leu Cys Gly Gly Val Leu Ile Asp Pro Gln Trp Val
 74 50 55 60
 77 Leu Thr Ala Ala His Cys Lys Lys Pro Asn Leu Gln Val Ile Leu Gly

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78	65	70	75	80
81	Lys His Asn Leu Arg Gln Thr Glu Thr Phe Gln Arg Gln Ile Ser Val			
82	85	90	95	
85	Asp Arg Thr Ile Val His Pro Arg Tyr Asn Pro Glu Thr His Asp Asn			
86	100	105	110	
89	Asp Ile Met Met Val His Leu Lys Asn Pro Val Lys Phe Ser Lys Lys			
90	115	120	125	
93	Ile Gln Pro Leu Pro Leu Lys Asn Asp Cys Ser Glu Glu Asn Pro Asn			
94	130	135	140	
97	Cys Gln Ile Leu Gly Trp Gly Lys Met Glu Asn Gly Asp Phe Pro Asp			
98	145	150	155	160
101	Thr Ile Gln Cys Ala Asp Val His Leu Val Pro Arg Glu Gln Cys Glu			
102	165	170	175	
105	Arg Ala Tyr Pro Gly Lys Ile Thr Gln Ser Met Val Cys Ala Gly Asp			
106	180	185	190	
109	Met Lys Glu Gly Asn Asp Ser Cys Gln Gly Asp Ser Gly Gly Pro Leu			
110	195	200	205	
113	Val Cys Gly Gly Arg Leu Arg Gly Leu Val Ser Trp Gly Asp Met Pro			
114	210	215	220	
117	Cys Gly Ser Lys Glu Lys Pro Gly Val Tyr Thr Asp Val Cys Thr His			
118	225	230	235	240
121	Ile Arg Trp Ile Gln Asn Ile Leu Arg Asn Lys Trp Leu			
122	245	250		
125	<210> SEQ ID NO: 3			
126	<211> LENGTH: 12			
127	<212> TYPE: DNA			
128	<213> ORGANISM: Artificial			
130	<220> FEATURE:			
131	<223> OTHER INFORMATION: Oligonucleotide adaptor for representational difference analysis			
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137	<210> SEQ ID NO: 4			
138	<211> LENGTH: 24			
139	<212> TYPE: DNA			
140	<213> ORGANISM: Artificial			
142	<220> FEATURE:			
143	<223> OTHER INFORMATION: Oligonucleotide adaptor for representational difference analysis			
145	<400> SEQUENCE: 4			
146	agcactctcc agcctctcac cgca	24		
149	<210> SEQ ID NO: 5			
150	<211> LENGTH: 12			
151	<212> TYPE: DNA			
152	<213> ORGANISM: Artificial			
154	<220> FEATURE:			
155	<223> OTHER INFORMATION: Oligonucleotide adaptor for representational difference analysis			
157	<400> SEQUENCE: 5			
158	gatctgttca tg	12		
161	<210> SEQ ID NO: 6			
162	<211> LENGTH: 24			

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Input Set : A:\31304-703.831.ST25.txt
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163 <212> TYPE: DNA
164 <213> ORGANISM: Artificial
166 <220> FEATURE:
167 <223> OTHER INFORMATION: Oligonucleotide adaptor forrepresentational difference analysis
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170 accgacgtcg actatccatg aaca 24
173 <210> SEQ ID NO: 7
174 <211> LENGTH: 12
175 <212> TYPE: DNA
176 <213> ORGANISM: Artificial
178 <220> FEATURE:
179 <223> OTHER INFORMATION: Oligonucleotide adaptor forrepresentational difference analysis
181 <400> SEQUENCE: 7 12
182 gatcttcccct cg
185 <210> SEQ ID NO: 8
186 <211> LENGTH: 24
187 <212> TYPE: DNA
188 <213> ORGANISM: Artificial
190 <220> FEATURE:
191 <223> OTHER INFORMATION: Oligonucleotide adaptor forrepresentational difference analysis
193 <400> SEQUENCE: 8
194 aggcaactgt gctatccgag ggaa 24

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Input Set : A:\31304-703.831.ST25.txt
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Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:3,4,5,6,7,8

VERIFICATION SUMMARY

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